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ANS 2 7 2004
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SEQUENCE LISTING

<110> Weber, Patricia C.
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 Madison, Vincent S.
 Wyss, Daniel
 Yao, Nanhua
 Liu, Dingjiang
 Gesell, Jennifer

<120> Hepatitis C Virus NS3 Helicase Fragments

<130> ID01152 US

<140> US 09/825,423

<141> 2001-04-03

<150> US 60/194,419

<151> 2000-04-04

<160> 16

<170> PatentIn Ver. 2.1

<210> 1

<211> 631

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Ile Ile Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly Glu
20 25 30

Val Gln Ile Val Ser Thr Ala Thr Gln Thr Phe Leu Ala Thr Cys Ile 35 40 45

Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr Ile 50 55 60

Ala Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp Gln 65 70 75 80

Asp Leu Val Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu Thr Pro 85 90 95

Cys Thr Cys Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala Asp 100 105 110

Val Ile Pro Val Arg Arg Gly Asp Ser Arg Gly Ser Leu Leu Ser 115 120 125

Pro Arg Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu Leu 130 135 140

Cys Pro Ala Gly His Ala Val Gly Leu Phe Arg Ala Ala Val Cys Thr 145 150 155 160

Arg Gly Val Thr Lys Ala Val Asp Phe Ile Pro Val Glu Asn Leu Glu

Thr Thr Met Arg Ser Pro Val Phe Thr Asp Asn Ser Ser Pro Pro Ala 185 Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr Gly Ser 200 Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly Tyr Lys Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala 230 235 Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr Gly Val Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Ile 280 Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu 310 Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Val Pro His Pro Asn Ile 325 330 Glu Glu Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr Gly Lys Ala Ile Pro Leu Glu Val Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val Ala Leu 375 Gly Ile Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Val Ile Pro Thr Asn Gly Asp Val Val Val Ala Thr Asp Ala Leu Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr Cys Val 425 Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu Thr 435 Thr Thr Leu Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg Gly Arg Thr Gly Arg Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro Gly Glu 475 Arg Pro Ser Gly Met Phe Asp Ser Ser Val Leu Cys Glu Cys Tyr Asp 490

Ala Gly Cys Ala Trp Tyr Glu Leu Thr Pro Ala Glu Thr Thr Val Arg 500 505 510

Leu Arg Ala Tyr Met Asn Thr Pro Gly Leu Pro Val Cys Gln Asp His 515 520 525

Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr His Ile Asp Ala 530 535 540

His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Phe Pro Tyr Leu 545 550 555 560

Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro Pro 575

Ser Trp Asp Gln Met Trp Lys Cys Leu Ile Arg Leu Lys Pro Thr Leu
580 585 590

His Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln Asn Glu 595 600 605

Val Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Thr Cys Met Ser 610 615 620

Ala Asp Leu Glu Val Val Thr 625 630

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<212> PRT

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<212> PRT

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Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr Gly Ser 20 25 30

Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly Tyr Lys 35 40 45

Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala 50 60

Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr Gly Val 65 70 75 80

Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys
85 90 95

Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Ile 100 105 110

Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile Gly
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Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu 130 135 140

Ala Thr Ala Thr 145

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<212> PRT

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Gly Ser His Met Gly Ser Val Thr Val Pro His Pro Asn Ile Glu Glu
1 5 10 15

Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr Gly Lys Ala Ile 20 25 30

Pro Leu Glu Val Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser 35 40 45

Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val Ala Leu Gly Ile 50 55 60

Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Val Ile Pro Thr 65 70 75 80

Asn Gly Asp Val Val Val Ala Thr Asp Ala Leu Met Thr Gly Phe 85 90 95

Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr Ser Asp Gly Lys
100 105 110

Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg Gly Arg Thr Gly Arg 115 120 125

Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro Gly Glu Arg 130 135 140

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<211> 288

<212> PRT

<213> Hepatitis C virus

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<211> 241

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Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr Gly Ser 20 25 30

Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly Tyr Lys 35 40 45

Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala
50 55 60

Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr Gly Val 65 70 75 80

Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys
85 90 95

Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Ile 100 105 110

Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile Gly
115 120 125

Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu 130 135 140

Ala Thr Ala Thr Pro Pro Gly Ser Gly Met Phe Asp Ser Ser Val Leu 145 150 155 160

Cys Glu Cys Tyr Asp Ala Gly Cys Ala Trp Tyr Glu Leu Thr Pro Ala 165 170 175

Glu Thr Thr Val Arg Leu Arg Ala Tyr Met Asn Thr Pro Gly Leu Pro 180 185 190

Val Cys Gln Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu 195 200 205

Thr His Ile Asp Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu 210 215 220

Asn Phe Pro Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala 225 230 235 240

Gln

<210> 7

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

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<400> 8
Arg Gly Ser Thr
<210> 9
<211> 4
<212> PRT
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<400> 9
Arg Gly Pro Gly
<210> 10
<211> 4
<212> PRT
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<223> Mutated Hepatitis C virus peptide
<400> 10
Ser Lys Gly Glu
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<212> PRT
<213> Artificial Sequence
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<400> 11
Glu Gln Gly Ala
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<400> 12
Arg Asn Asn Gln
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Ala Asp Gly Ser
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<211> 4
<212> PRT
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Cys Asp Gly Leu
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<211> 33
<212> DNA
<213> Artificial Sequence
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<211> 33
<212> DNA
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33